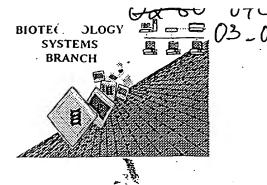
BEST AVAILABLE COPY

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09/778,97/
Source:	OIPE
Date Processed by STIC:	3-1-01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 c-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 c-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

Does Not Comply Corrected Diskette Needed

Add second bracket to numerical headers. Delete Delete

SEQUENCE LISTING <110/Shaughnessy, John D. 4120/Evi27 Gene Sequence and
Protein Encoded Thereby <130/D6138PCT <140/<141/2001-02-02 <150/US 60/180,374
<151 2000-02-04 <160 9 <210 1 <211 1827 <212 DNA, <213 Homo sapiens <220 <221

prim_transcript <223 cDNA of human Evi27 <400 1 cggcgatgtc gctcgtgctg ataagcctgg
ccgcgctgtg caggagcgcc 50 gtaccccgag agccgaccgt tcaatgtggc tctgaaactg ggccatctcc Add spaces between headers agagtggatg ctacaacatg atctaatccc cggagacttg agggacctcc 150 gagtagaacc tgttacaact agtgttgcaa caggggacta ttcaattttg 200 atgaatgtaa gctgggtact ccgggcagat gccagcatcc gcttgttgaa ggccaccaag atttgtgtga cgggcaaaag caacttccag tcctacagct 300 gtgtgaggtg caattacaca agactcagac cagaccetet 350 ggtggtaaat ggacatttte etatategge tteeetgtag agetgaacae agtictattte attggggccc ataatattcc taatgcaaat atgaatgaag 450 atggcccttc catgtctgtg caccaggetg cetagaceae 500 Lataatgaaat ataaaaaaaa gtgtgteaag geeggaagee tgtgggatee gaacatcact gcttgtaaga agaatgagga gacagtagaa gtgaacttca 600 caaccactcc cctgggaaac agatacatgg ctcttatcca acacagcact 650 atcatcgggt tttctcaggt gtttgagcca caccagaaga aacaaacgcg 700 agetteagtg gtgattecag tgactgggga tagtgaaggt getacggtge 750 agetgactee atattttcct acttgtggca gcgactgcat ccgacataaa 800 ggaacagttg tgctctgccc acaaacaggc gtccctttcc ctctggataa 850 caacaaaagc aagccgggag gctggctgcc tctcctcctg ctgtctctgc 900 tggtggccac gtggcaggga tctatctaat gtggaggcac 950 gaaaggatca agaagacttc cttttctacc accacactac tgcccccat 1000 taaggttett gtggtttace catetgaaat atgttteeat cacacaattt 1050 gttaetteae tgaatttctt caaaaccatt gcagaagtga ggtcatcctt 1100 gaaaagtggc agaaaagaa aatagcagag atgggtccag tgcagtggct 1150 tgccactcaa aagaaggcag cagacaaagt cgtcttcctt ctttccaatg 1200 acgtcaacag tgtgtgcgat ggtacctgtg gcaagagcga gggcagtccc 1250 agtgagaact ctcaagacct cttccccctt gcctttaacc ttttctgcag 1300 tgatctaaga agccagattc atctgcacaa atacgtggtg gtctacttta 1350 gagagattga tacaaaagac gattacaatg ctctcagtgt ctgccccaag 1400 taccacttca tgaaggatgc cactgctttc tgtgcagaac ttctccatgt 1450 caagcagcag gtgtcagcag gaaaaagatc acaagcctgc cacgatggct 1500 gctgctcctt catgagaagc aagagacctt aaaggcttcc 1550 tatcccacca attacaggga aaaaacgtgt gatgatcctg aagcttacta 1600 tgcagcctac aaacagcctt agtaattaaa acattttata ccaataaaat 1650 tttcaaatat tactaactaa tgtagcatta actaacgatt ggaaactaca 1700 tttacaactt caaagctgtt ttatacatag aaatcaatta cagctttaat 1750 tgaaaactgt

aaccattttg ataatgcaac aataaagcat cttccaaaaa 1800 aaaaaaaaaa aaaaaaaa aaaaaaaa

Does Not Comply
Corrected Diskette Needed

Format errors.

Format errors.

Insert "hard returns"

Undicate.

Where orrows indicated in

These errors are indicated in

all of the sequences